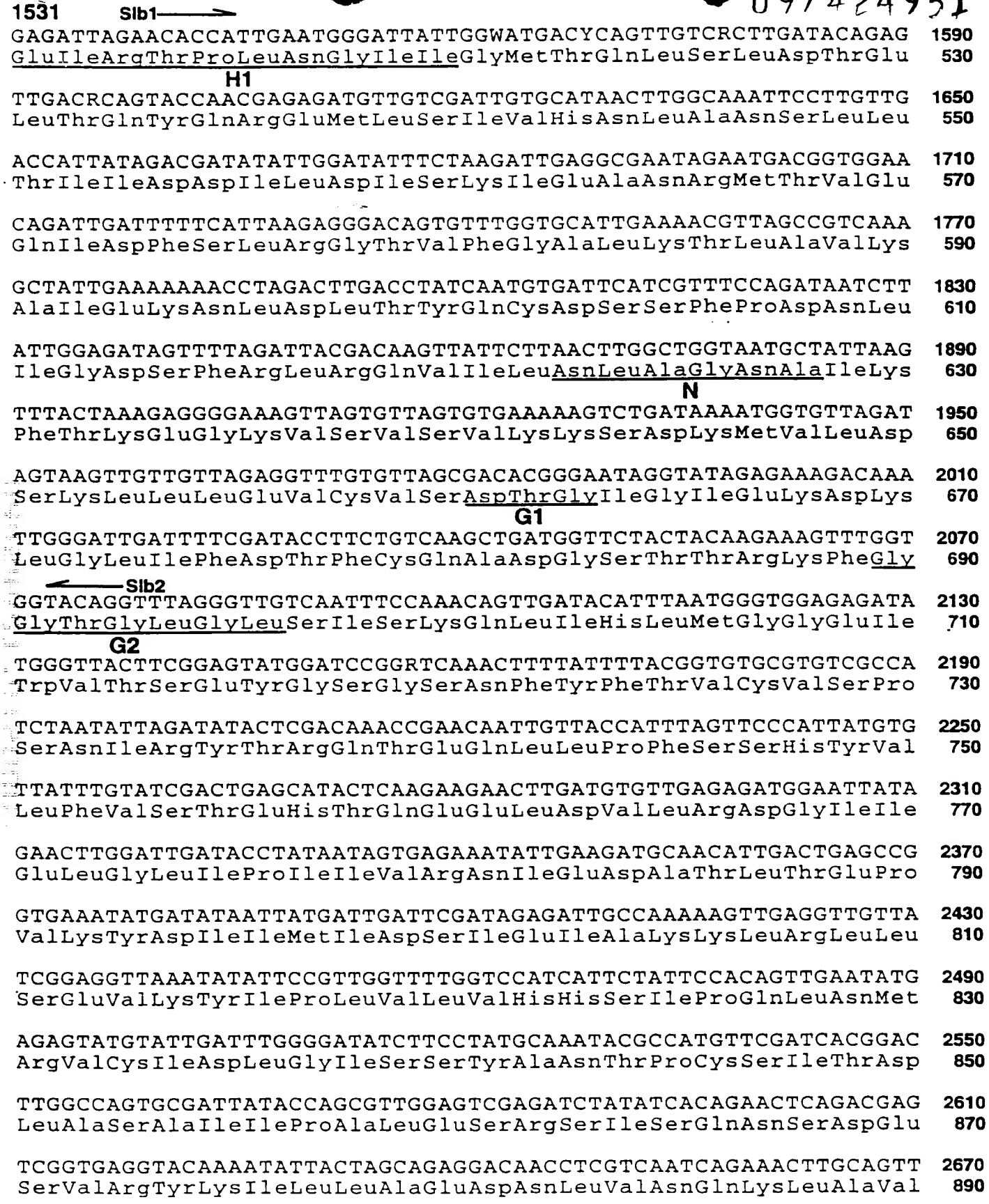
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LeuThrGlnTyrGlnArgGluMetLeuSerIleValHisAsnLeuAlaAsnSerLeuLeu 550
ACCATTATAGACGATATATTGGATATTTCTAAGATTGAGGCGAATAGAATGACGGTGGAA 1710
ThrIleIleAspAspIleLeuAspIleSerLysIleGluAlaAsnArgMetThrValGlu 570
CAGATTGATTTTTCATTAAGAGGGACAGTGTTTGGTGCATTGAAAACGTTAGCCGTCAAA 1770
GlnIleAspPheSerLeuArgGlyThrValPheGlyAlaLeuLysThrLeuAlaValLys 590
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AlaIleGluLysAsnLeuAspLeuThrTyrGlnCysAspSerSerPheProAspAsnLeu 610
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IleGlyAspSerPheArgLeuArgGlnValIleLeuAsnLeuAlaGlyAsnAlaIleLys 630
N
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PheThrLysGluGlyLysValSerValSerValLysLysSerAspLysMetValLeuAsp 650
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SerLysLeuLeuLeuGluValCysValSerAspThrGlyIleGlyIleGluLysAspLys 670
G1
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LeuGlyLeuIlePheAspThrPheCysGlnAlaAspGlySerThrThrArgLysPheGly 690
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GlyThrGlyLeuGlyLeuSerIleSerLysGlnLeuIleHisLeuMetGlyGlyGluIle 710
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LeuAlaSerAlaIleIleProAlaLeuGluSerArgSerIleSerGlnAsnSerAspGlu 870
TCGGTGAGGTACAAAATATTACTAGCAGAGGACAACCTCGTCAATCAGAAACTTGCAGTT 2670
SerValArgTyrLysIleLeuLeuAlaGluAspAsnLeuValAsnGlnLysLeuAlaVal 890

Fig. 1a

```
AGGATATTAGAAAAGCAAGGGCATCTGGTGGAAGTAGTTGAGAACGGACTCGAGGCGTAC  2730
ArgIleLeuGluLysGlnGlyHisLeuValGluValValGluAsnGlyLeuGluAlaTyr   910

                                                  <--- Slb3
GAAGCGATTAAGAGGAATAAATATGATGTGGTGTTGATGGATGTGCAAATGCCT         2784
GluAlaIleLysArgAsnLysTyrAspValValLeuMetAspValGlnMetPro          928
                                    D
```

Fig. 1b

```
ATGAACCCCACTAAAAAACCTCGGTTATCACCAATGCAGCCCTCTGTTTTTGAAATACTC   60
MetAsnProThrLysLysProArgLeuSerProMetGlnProSerValPheGluIleLeu   20

AACGACCCTGAGCTTTATAGTCAGCACTGTCATAGCCTTAGGGAAACACTTCTTGATCAT  120
AsnAspProGluLeuTyrSerGlnHisCysHisSerLeuArgGluThrLeuLeuAspHis   40

TTCAACCATCAAGCTACACTTATCGACACTTATGAACATGAACTAGAAAAATCCAAAAAT  180
PheAsnHisGlnAlaThrLeuIleAspThrTyrGluHisGluLeuGluLysSerLysAsn   60

GCCAACAAAGCGTCCCAACAAGCACTTAGTGAAATAGGTACAGTTGTTATATCTGTTGCC  240
AlaAsnLysAlaSerGlnGlnAlaLeuSerGluIleGlyThrValValIleSerValAla   80

ATGGGAGACTTGTCGAAAAAAGTTGAGATTCACACAGTAGAAAATGACCCTGAGATTTTA  300
MetGlyAspLeuSerLysLysValGluIleHisThrValGluAsnAspProGluIleLeu  100

AAAGTCAAAATCACCATCAACACCATGATGGATCAATTACAGACATTTGCTAATGAGGTT  360
LysValLysIleThrIleAsnThrMetMetAspGlnLeuGlnThrPheAlaAsnGluVal  120

ACAAAAGTCGCCACCGAAGTCGCAAATGGTGAACTAGGTGGACAAGCGAAAAATGATGGA  420
ThrLysValAlaThrGluValAlaAsnGlyGluLeuGlyGlyGlnAlaLysAsnAspGly  140

TCTGTTGGTATTTGGAGATCACTTACAGACAATGTTAATATTATGGCTCTTAATTTAACT  480
SerValGlyIleTrpArgSerLeuThrAspAsnValAsnIleMetAlaLeuAsnLeuThr  160

AACCAAGTGCGAGAAATTGCTGATGTCACACGTGCTGTTGCCAAGGGGGACTTGTCACGT  540
AsnGlnValArgGluIleAlaAspValThrArgAlaValAlaLysGlyAspLeuSerArg  180

AAAATTAATGTACACGCCCAGGGTGAAATCCTTCAACTTCAACGTACAATAAACACCATG  600
LysIleAsnValHisAlaGlnGlyGluIleLeuGlnLeuGlnArgThrIleAsnThrMet  200

GTGGATCAGTTACGAACGTTTGCATTCGAAGTATCTAAAGTTGCTAGAGATGTTGGTGTG  660
ValAspGlnLeuArgThrPheAlaPheGluValSerLysValAlaArgAspValGlyVal  220

CTTGGTATATTAGGAGGACAAGCGTTGATTGAAAATGTTGAAGGTATTTGGGAAGAGTTG  720
LeuGlyIleLeuGlyGlyGlnAlaLeuIleGluAsnValGluGlyIleTrpGluGluLeu  240

ACTGATAATGTCAATGCCATGGCTCTTAATTTGACTACACAAGTGAGAAATATTGCCAAT  780
ThrAspAsnValAsnAlaMetAlaLeuAsnLeuThrThrGlnValArgAsnIleAlaAsn  260

GTCACCACTGCCGTTGCCAAGGGGGATTTGTCGAAAAAAGTCACTGCTGATTGTAAGGGA  840
ValThrThrAlaValAlaLysGlyAspLeuSerLysLysValThrAlaAspCysLysGly  280

GAAATYCTTGATTTGAAACTTACTATTAATCAAATGGTGGACCGATTACAGAATTTTGCT  900
GluIleLeuAspLeuLysLeuThrIleAsnGlnMetValAspArgLeuGlnAsnPheAla  300

CTTGCGGTGACGACATTGTCGAGAGAGGTTGGTACTTTGGGTATTTTGGGTGGACAAGCT  960
LeuAlaValThrThrLeuSerArgGluValGlyThrLeuGlyIleLeuGlyGlyGlnAla  320

AACGTACAGGATGTTGAAGGTGCTTGGAAACAGGTTACAGAAAATGTCAACCTAATGGCT 1020
AsnValGlnAspValGluGlyAlaTrpLysGlnValThrGluAsnValAsnLeuMetAla  340

ACTAATTTAACTAACCAAGTGAGATCTATTGCTACAGTTACTACTGCAGTTGCGCATGGT 1080
ThrAsnLeuThrAsnGlnValArgSerIleAlaThrValThrThrAlaValAlaHisGly  360

GATTTGTCGCAAAAGATTGATGGTCATCCCAAAGGAGAGATTTTACAATTGAAAAATACA 1140
AspLeuSerGlnLysIleAspGlyHisProLysGlyGluIleLeuGlnLeuLysAsnThr  380
```

Fig. 2a

```
ATCAACAAGATGGTGGACTC●GCAGTTGTTTGCATCAGAAGTG●GAAAGTGGCACAA 1200
IleAsnLysMetValAspSerLeuGlnLeuPheAlaSerGluValSerLysValAlaGln 400

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AspValGlyIleAsnGlyLysLeuGlyIleGlnAlaGlnValSerAspValAspGlyLeu 420

TGGAAGGAGATTACGTCTAATGTAAATACCATGGCTTCAAATTTAACTTCGCAAGTGAGA 1320
TrpLysGluIleThrSerAsnValAsnThrMetAlaSerAsnLeuThrSerGlnValArg 440

GCTTTTGCACAGATTACTGCTGCTGCTACTGATGGGGATTTCACTAGATTTATTACTGTT 1380
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GluAlaLeuGlyGluMetAspAlaLeuLysThrLysIleAsnGlnMetValPheAsnLeu 480

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ArgGluSerLeuGlnArgAsnThrAlaAlaArgGluAlaAlaGluLeuAlaAsnSerAla 500

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LysSerGluPheLeuAlaAsnMetSerHisGluIleArgThrProLeuAsnGlyIleIle 520
                                        H1
GGWATGACYCAGTTGTCRCTTGATACAGAGTTGACRCAGTACCAACGAGAGATGTTGTCG 1620
GlyMetThrGlnLeuSerLeuAspThrGluLeuThrGlnTyrGlnArgGluMetLeuSer 540

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IleValHisAsnLeuAlaAsnSerLeuLeuThrIleIleAspAspIleLeuAspIleSer 560

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PheGlyAlaLeuLysThrLeuAlaValLysAlaIleGluLysAsnLeuAspLeuThrTyr 600

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IleLeuAsnLeuAlaGlyAsnAlaIleLysPheThrLysGluGlyLysValSerValSer 640
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AspThrGlyIleGlyIleGluLysAspLysLeuGlyLeuIlePheAspThrPheCysGln 680
G1
GCTGATGGTTCTACTACAAGAAAGTTTGGTGGTACAGGTTTAGGGTTGTCAATTTCCAAA 2100
AlaAspGlySerThrThrArgLysPheGlyGlyThrGlyLeuGlyLeuSerIleSerLys 700
                                        G2
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AsnPheTyrPheThrValCysValSerProSerAsnIleArgTyrThrArgGlnThrGlu 740

CAATTGTTACCATTTAGTTCCCATTATGTGTTATTTGTATCGACTGAGCATACTCAAGAA 2280
GlnLeuLeuProPheSerSerHisTyrValLeuPheValSerThrGluHisThrGlnGlu 760

GAACTTGATGTGTTGAGAGATGGAATTATAGAACTTGGATTGATACCTATAATAGTGAGA 2340
GluLeuAspValLeuArgAspGlyIleIleGluLeuGlyLeuIleProIleIleValArg 780
```

Fig. 2b

```
AATATTGAAGATGCAACATTGACTGAGCCGGTGAAATATGATATAATTATGATTGATTCG 2400
AsnIleGluAspAlaThrLeuThrGluProValLysTyrAspIleIleMetIleAspSer  800

ATAGAGATTGCCAAAAAGTTGAGGTTGTTATCGGAGGTTAAATATATTCCGTTGGTTTTG 2460
IleGluIleAlaLysLysLeuArgLeuLeuSerGluValLysTyrIleProLeuValLeu  820

GTCCATCATTCTATTCCACAGTTGAATATGAGAGTATGTATTGATTTGGGGATATCTTCC 2520
ValHisHisSerIleProGlnLeuAsnMetArgValCysIleAspLeuGlyIleSerSer  840

TATGCAAATACGCCATGTTCGATCACGGACTTGGCCAGTGCGATTATACCAGCGTTGGAG 2580
TyrAlaAsnThrProCysSerIleThrAspLeuAlaSerAlaIleIleProAlaLeuGlu  860

TCGAGATCTATATCACAGAACTCAGACGAGTCGGTGAGGTACAAAATATTACTAGCAGAG 2640
SerArgSerIleSerGlnAsnSerAspGluSerValArgTyrLysIleLeuLeuAlaGlu  880

GACAACCTCGTCAATCAGAAACTTGCAGTTAGGATATTAGAAAAGCAAGGGCATCTGGTG 2700
AspAsnLeuValAsnGlnLysLeuAlaValArgIleLeuGluLysGlnGlyHisLeuVal  900

GAAGTAGTTGAGAACGGACTCGAGGCGTACGAAGCGATTAAGAGGAATAAATATGATGTG 2760
GluValValGluAsnGlyLeuGluAlaTyrGluAlaIleLysArgAsnLysTyrAspVal  920

GTGTTGATGGATGTGCAAATGCCTGTAATGGGTGGGTTTGAAGCTACGGAGAAGATTCGA 2820
ValLeuMetAspValGlnMetProValMetGlyGlyPheGluAlaThrGluLysIleArg  940
          D
CAATGGGAGAAAAAGTCTAACCCAATTGACTCGTTGACGTTTAGGACTCCAATTATTGCC 2880
GlnTrpGluLysLysSerAsnProIleAspSerLeuThrPheArgThrProIleIleAla  960

CTCACTGCACACGCCATGTTAGGTGATAGAGAAAAGTCATTGGCCAAGGGGATGGACGAT 2940
LeuThrAlaHisAlaMetLeuGlyAspArgGluLysSerLeuAlaLysGlyMetAspAsp  980

TATGTGAGTAAGCCATTGAAGCCGAAATTGTTAATGCAGACGATAAACAAGTGTATTCAT 3000
TyrValSerLysProLeuLysProLysLeuLeuMetGlnThrIleAsnLysCysIleHis 1000
                                          H2
AATATTAACCAGTTGAAAGAATTGTCGAGAAATAGTAGGGGTAGCGATTTTGCAAAGAAG 3060
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MetThrArgAsnThrProGlySerThrThrArgGlnGlySerAspGluGlySerValLys 1040

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AspMetIleGlyAspThrProArgGlnGlySerValGluGlyGlyGlyThrSerSerArg 1060

CCAGTACAGAGAAGGTCTGCCAGGGAGGGGTCGATCACTACAATTAGTGAACAAATCGAC 3240
ProValGlnArgArgSerAlaArgGluGlySerIleThrThrIleSerGluGlnIleAsp 1080

CGTTAG                                                         3246
Arg***                                                         1082
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Fig. 2c

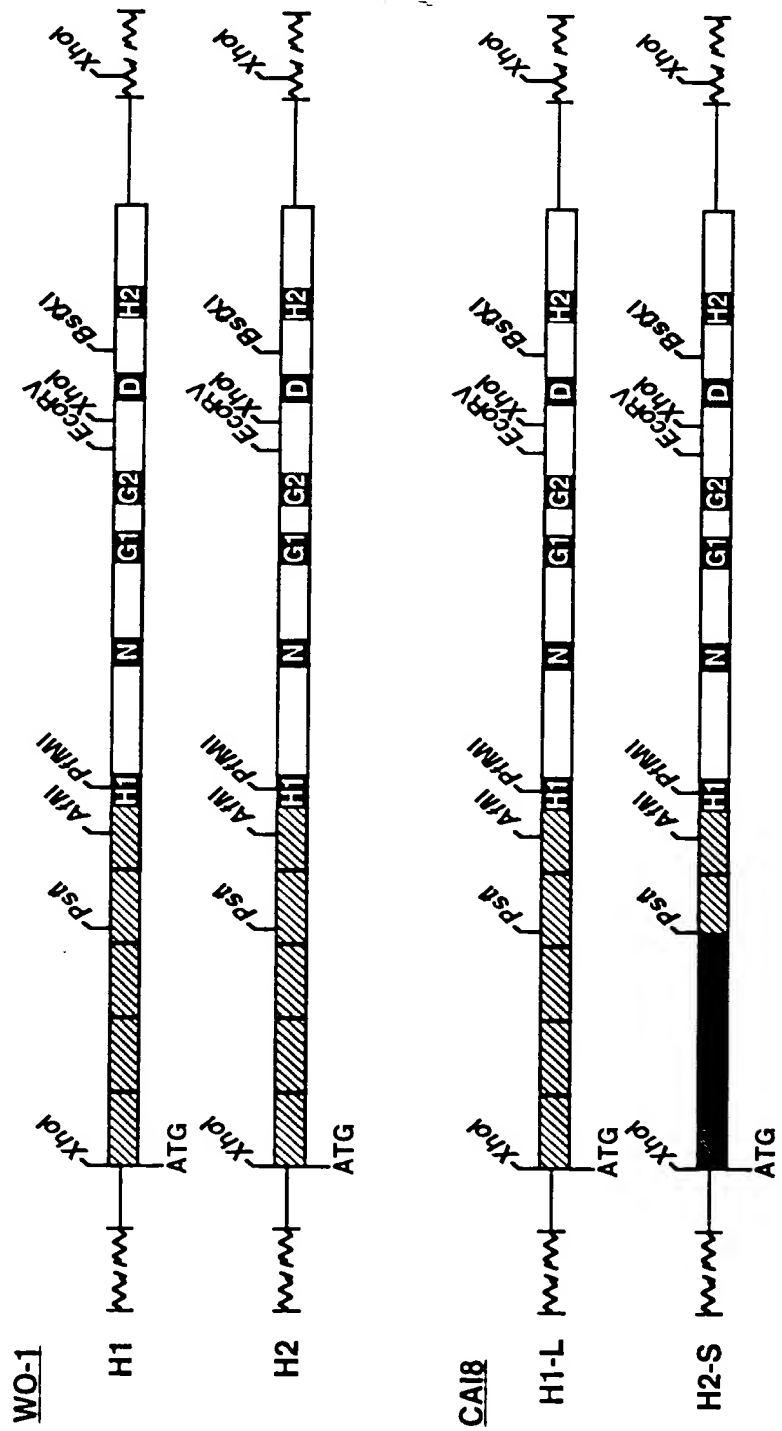
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Fig. 3

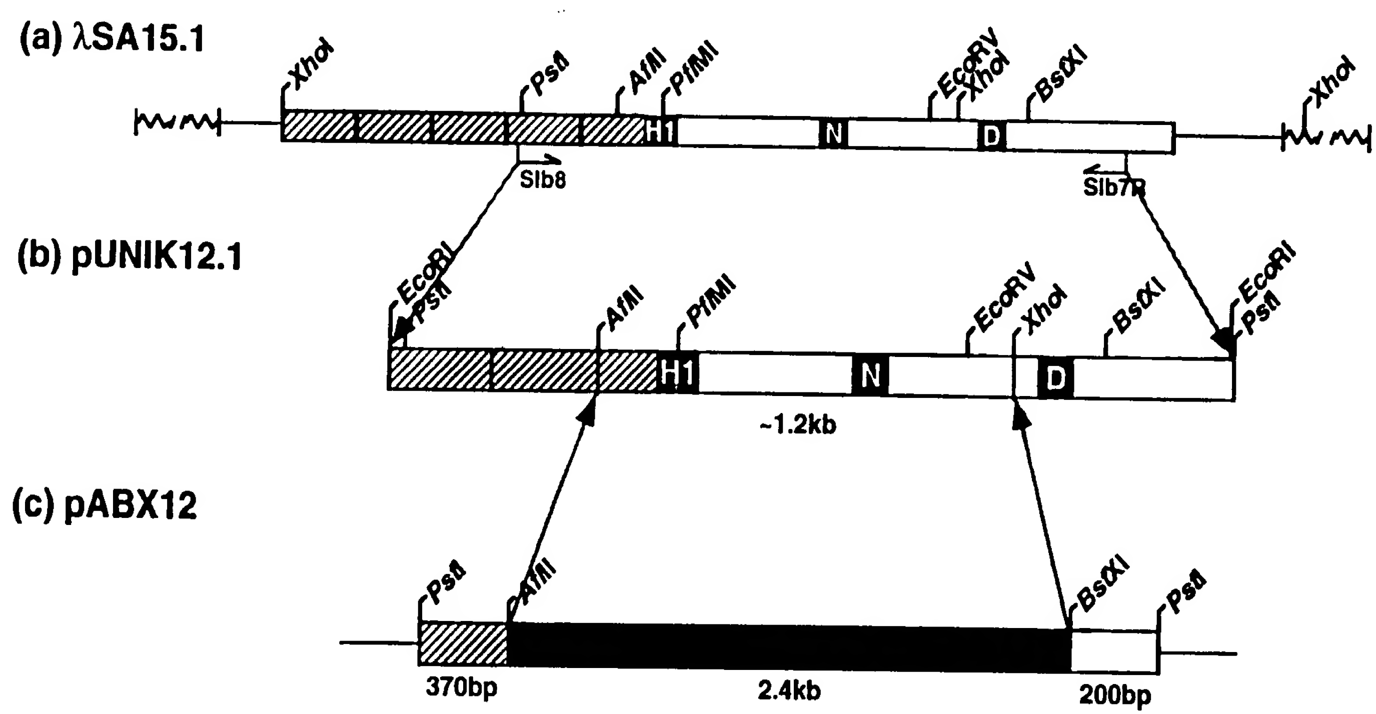
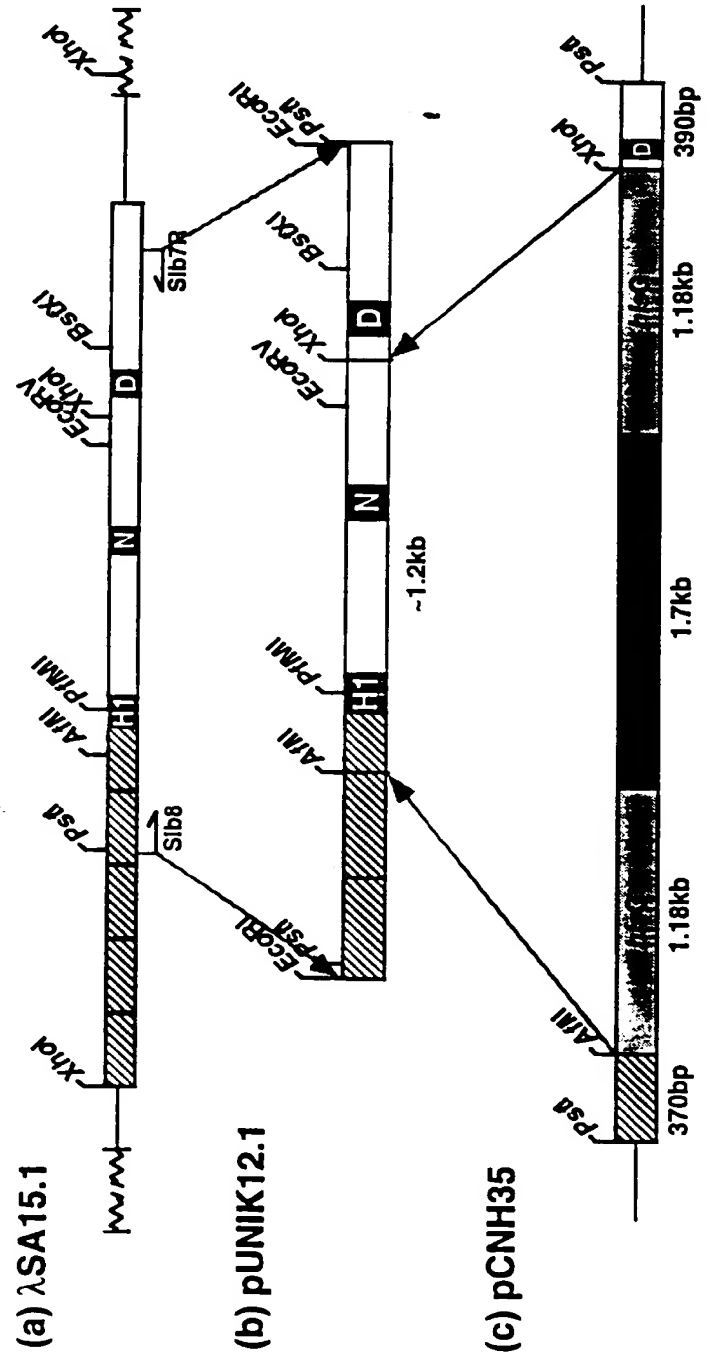


Fig. 4

09/424951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370bp
PstI

Fig. 5